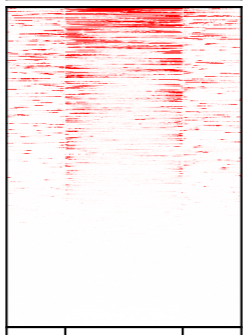
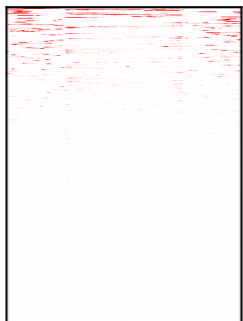


H3K79me2_K562_ENCFF334HSS

498unaffected_hg38_GenCode44bedregulated_hg38_GenCode44bed



-20.000 20.000
gene distance (bp)